

# FIG. 1A-1

GGAGGTATAGGAGCTCTCTTCGATTTTAGCAAAACCAGGAGTCCGAAGATCTAAGGAGAGCTGGGGTTTGACTCC

SacI

85 95 105 115 125 135 145

GAGAGCTCGAGCAGTCCCAAGACCTGGTCTTGACTCAGGAGTTAGACTCCACTCAGAGGCTGACTGTCTCCAGG

SacI PflMI

XhoI TthlIII

160 170 180 190 200 210 220

GTCTACACCTCTAAGGGCGACACTGGGCTCAAGCAGACTGCCGTTTCTATATGGGATGAGCCTTCACAGGGCAG

235 245 255 265 275 285 295

CCAGTTGGGATGGGTTGAGGTTTGGCTGTAGACATCAGAAACCCAAAGTCAAATGCGCTTCAACCAGTAGAAAATT

310 320 330 340 350 360 370

CACCAGCCCGCAGAGCTAAGGTTGGGTGGACATTAGGGTTGGTTGATCCAGGAGCTCAACAGTGTCTCTGAGCC

SacI

385 395 405 415 425 435 445

CCAGCTCCTTCTGCCCCCACCACCATCTTCAGTGTCTCTCAAGGCCACAGCTGTAGTTGGCCAGGGGG

PvuII BglI

BglI

460 470 480 490 500 510 520

GCTTCATTATTTTGGCTCCTGGGCAGTAGGAGGAAGAGAAATGAATGTCTCTCCATGGGTCTTCTTAGGAATGT

NcoI

535 545 555 565 575 585 595

GGGAACCTTTTCCAGAACTCTATGTCTTTTAGTTTGTGGTCACTTGCCCTTCTCTGAACCACTTCTCTGAC

610 620 630 640 650 660 670

TCCTGGACAGGATGTGCACCTGATGAGCTTAGCTTTGGGGATCTAATAGTGACTTTACAAAGCCCTCTTTGAGAAGG

ApalI EspI

685 695 705 715 725 735 745

TGACATTGGAACCAAGGCTTGAGCAGACACAACAAGATTGCAGGGAGGGGCATTGCAGGTGGAGGAAACGGCAC

BspMI-

760 770 780 790 800 810 820

ATGCAAGAGCCCTGCGTGGGAGTGAGCTTGGTGTGTTGGTCAATCAGTTGTCTCAGAGCACACCGGGCCCTGTCTAGCA

ApalI

EcoO

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APPROVED BY CLASS SUBCLAS

530 326

5258494

Fig 1-49

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# FIG. 1A-2

APPROVED	0.6 FIG. 1A-2
BY	CLASS
DATE	SUBCLASS
530	326

835 845 855 865 875 885 895  
 GGCACAGCCTGGGCCTGCTCTGAGTATGACAGAGAGCCCTGGGAAGTTGTAGGTGGAGGAAGACAGAGTTCATGA  
 910 920 930 940 950 960 970  
 CTAGGAAAAAGCAATCCCTCTGTTGTGGGTGGAAGGTTGCAGTGTGTGTGAGAGAGAGACAAGACAGAC  
 985 995 1005 1015 1025 1035 1045  
 AGACAGACACTTCTCAATGTTTACAAGTGCTCAGGCCCTGACCCCGAATGCTTCCAAATTTACGTAGTTCTTGAA  
 EcoO BsmI+ SnaBI  
 1060 1070 1080 1090 1100 1110 1120  
 ACCCCCTGTATCATTTTCACTACTCAAGAAACCTCGGAGTGTCTTCTCTGAAAGTTCATCAGGTTTGTACTC  
 1135 1145 1155 1165 1175 1185 1195  
 TCTGCTGTCTCATTTCTTCTTGTGCTGGTGATGGTTGCTTGTCCCGCCCTGTCCCGCATCCTCTTGTCCC  
 EcoO  
 1210 1220 1230 1240 1250 1260 1270  
 CTGCAGAGGGATGAGTGTGTGGGGCCTCACGAGTTGAGTTGTTTCATAAGCAGATCTCTTTGAGCAGGGCGCCT  
 PstI EcoO BglII NarI Ps  
 1285 1295 1305 1315 1325 1335 1345  
 GCAGTGGCCTTGTGTGAGGCTGGAGGGGTTTCGATTCCCTTATGGAATCCAGGCAGATGTAGCATTTAAACAACA  
 tI DraI  
 1360 1370 1380 1390 1400 1410 1420  
 CACGTGTATAAAGAAACCAGTGTCCGCAGAAAGTTCCAGAAAGTATTATGGGATAAGACTACATGAGAGAGGAA  
 1435 1445 1455 1465 1475 1485 1495  
 TGGGGCAATTGGCACCTCCCTTAGTAGGGCCTTTGCTGGGGGTAGAAATGAGTTTAAAGGCAGGTTAGACCCCTCGA  
 EcoO BspMI-  
 1510 1520 1530 1540 1550 1560 1570  
 ACTGGCTTTTGAATCGGGAAATTTACCCCCCAGCCGTTCTGTGCTTCAATTGCTGTTACATCACTGCCTAAGATG  
 1585 1595 1605 1615 1625 1635 1645  
 GAGGAACCTTTGATGTGTGTGTTTCTTCTCCCTCACTGGGCTCTGCTTCTTCACTTCTTGTCAATGCAGAGAA  
 1660 1670 1680 1690 1700 1710 1720  
 CAGCAGCAGGCCACAGAGGCAGGCCCTTGTAAAGACGAGCTGTATGTCAGCTTCCGAGACCTGGGCTGGCAGG  
 StuI BspMI  
 1735 1745 1755 1765 1775 1785 1795  
 TAAGGGGCTGGCTGGTCTGTCTTGGGTGTGGGCCCTCTGTGGCGTGGGCTCCACAGGCAGCGGGTGTGTGCTCA  
 ApaI  
 EcoO

556

1810 1820 1830 1840 1850 1860 1870  
GTCTTGTTCTCATCTCTGCCAGTTAAGACTCCAGTATCAAGTGGCTCGCTAGGGAAGGTACTTGGCTAAGGA  
1885 1895 1905 1915 1925 1935 1945  
TACAGG.....(APPROX. 1000 BASES).....GGAGCCAGCATGGGTGATGCCATTATGA  
1960 1970 1980 1990 2000 2010 2020  
GTTATTAGCCCTCTCTGGCAGGTGGGCAACCGAGGCATGGAGGTTGTTAAGGTGAACCTGCCAGTGTGTGACCA

BglI BspMI-  
2035 2045 2055 2065 2075 2085 2095  
CCTAGTGGGTAGAGCTGATGATTGCCTCACACCGAGCTCCTTCCTGTGCCGCTTCTGTCCAGAAGACACAGC  
aIII  
MI  
SacI  
N

# FIG. 1A-3

2110 2120 2130 2140 2150 2160 2170  
CATGGATGTCCATTTTAGGATCAGCCAGCCCGCTTGTCTTCATTTTATTTTATGTTTATTAGAAATGGG  
col

2185 2195 2205 2215 2225 2235 2245  
GTCTTGCTCTGCACCCAGGCTGGGTGCAGTGGTGTGATCATAGCTCACCGCAGCTTTGACGCCGCTCTTCCCACCT  
TthIII

2260 2270 2280 2290 2300 2310 2320  
CAGTCTACTAAGCTTGACTATAGGCCAAGACTATAGAGTGTGCTCTTCTTCCATTCTTTTGGACCATGAGAGG  
HindIII BstXI

2335 2345 2355 2365 2375 2385 2395  
CCACCCATGTTTCCTGCCCCCTGCTGGGCCCTGCTGCTCAGAAAGGCATGGTCTGAGGCTTTCACCTTGGTCTGTGAG  
ApaI  
EcoO

2410 2420 2430 2440 2450 2460 2470  
CCTTCGTGGTGGTTCTTTACAGCATGGGGTTGGATGCTGTGCTCAGGCTTCTGCATGGTTTCCCACACTCTCTT  
2485 2495 2505 2515 2525 2535 2545  
CTCCTCCTCAGGACTGGATCATCGCGCCTGAAGGCTACGCGGCTACTGTGAGGGGGAGTGTGCCCTTCCCTC

MstII BssHII  
2560 2570 2580 2590 2600 2610 2620  
TGAACTCCTACATGAACGCCACCAACCGCCATCGTGCAGACGCTGGTGGTGTACGCCCATCTTGGGGTGTGG  
Bs

2635 2645 2655 2665 2675 2685 2695  
TCACCTGGCCGGCAGGCTGCGGGGCCACAGATCCTGCTGCCTCCAAGCTGGGGCCTGAGTAGATGTCAGCCC  
tEII BglI  
EcoO

25

## FIG. 1A-4

2710 2720 2730 2740 2750 2760 2770  
ATTGCCATGTCATGACTTTTGGGGGGCCCCCTTGGCGCGTTAAAAAAATCAAAAATTGTACTTTATGACTGGTTT

2785 2795 2805 2815 2825 2835 2845  
GGTATAAGAGGAGTATAATCTTCGACCCCTGGAGTTCAATTTCTCCTAATTTTAAAGTAACTAAAAGTTGT

2860 2870 2880 2890 2900 2910 2920  
ATGGGCTCCTTTGAGGATGCTTGTAGTATTGTGGTGCTGGTTACGGTGCCCTAAGAGCACTGGGGCCCCCTGCTTCA

2935 2945 2955 2965 2975 2985 2995  
TTTTCCAGTAGAGGAAACAGGTAAACAGATGAGAAATTTTCAGTGAGGGGCACAGTGATCAGAAAGGGGCCAGCAG

3010 3020 3030 3040 3050 3060 3070  
GATAATGGGATGGAGAGATGAGTGGGGACCCATGGGCCATTTCAAGTTAAATTTTCAGTCGGGTCCACCAGGAAGAT

3085 3095 3105 3115 3125 3135 3145  
TCCATGTGATAATGAGATTAAACGTGCCCGCCAGTCACGGCGACACTCAGTAGGTGTTATTCTCTGCTCTGCCAACAGCA

3160 3170 3180 3190 3200 3210 3220  
ACCATAGTTGATAAGAGCTGTTAGGATTTTGTCTTTTGTCTTAGAATCCAAGGTTCAAGGACCTTGGTTATGTA

3235 3245 3255 3265 3275 3285 3295  
GCTCCCTGTCATGAACATCATCTGAGCCTTTCTCTGCCTACTGATCATCCACCCCTGCCCTTGAATGCTTCTAGTGAC

3310 3320 3330 3340 3350 3360 3370  
AGAGAGCTCACTACCAGGACTACTCCCTCCTTTTCATTTAGTAATCTGCCCTCCTTCTTTTCTTGTCCTGTCTCTGT

3385 3395 3405 3415 3425 3435 3445  
GTGTTAAGTCCCTGGAGAAAAATCTCATCTATCCCTTTTCATTTGATTCTGCTCTTTTGAGGGCAGGGGTTTTTGT

3460 3470 3480 3490 3500 3510 3520  
CTTTGTTGTTTTTTTAAAGTGTGGTTTTTCCAAAGCCCTTGCTCCCTCCTCAATTGAAACTTCAAAGCCCTCAT

3535 3545 3555 3565 3575 3585 3595  
TGGGATTGAAGGTCCTTAGGCTGGAAACAGAGAGTCTCCCTCCCAACCTGTTCCCTGGCCCTGGATGTGCTGTGCTG

ECOOmStII

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APPROVED	0.C. FIG. 1A-4
BY	CLASS
DRAFTSMAN	SUBCLASS
520	726

# FIG. 1A-5

3610 3620 3630 3640 3650 3660 3670  
 TGCCAGTATCCCTGGAAAGTGCCAGGCATGTCTCCCGGCTGCCAGGGACACATCTCTATCTCTCTCAACCC  
 3685 3695 3705 3715 3725 3735 3745  
 CTGCCTTCATGGCCCATGGAACAGGAGTGCCATGCCCCTGTGTGCACCTACTTCCATCAGTATTCACCAAGAT  
 BglI NcoI ApaI BglI  
 3760 3770 3780 3790 3800 3810 3820  
 CTGCAGGATCAAAAGTGAATTCTCCAGGGATTGTGAAATGATGCGATTGTGGTCATGTTTAAAGGGGCAACTGT  
 I EcoRI DraI  
 3835 3845 3855 3865 3875 3885 3895  
 CTTCTAGAGAGTCCCTGATGAAATGCTTCCAGAGGAAATGAGCTGATGGCTGGAATTTGCTTTAAATCATTTCAAG  
 XbaI  
 3910 3920 3930 3940 3950 3960 3970  
 GTGGAGCAGGTGGGGAAGGTATGGATGTGTAAAGATTGTCCATCATATAAATGTGTAAAGCATGCT  
 BspMI- SphI  
 3985 3995 4005 4015 4025 4035 4045  
 GGCCTATGTCAGCAGTCACAGCCTGGAGGTGGTAACAGAGTGCCAGTCACTGATGCTCAAGCCTGGCACCCTACAG  
 4060 4070 4080 4090 4100 4110 4120  
 TTGCTGGAACCCAGAAAGTTTCACGTTGAAAACAACAGGACAGTGGAATCTCTGGCCCCTGTCTTGAACACCGTGGC  
 4135 4145 4155 4165 4175 4185 4195  
 AGATCTGCTAACACTGATCTTGGTTGGCTGCCGTCAGCTTAGGTTGAGTGGCGGTCTTCCCTTAGTTTGCTTAGT  
 BglII  
 4210 4220 4230 4240 4250 4260 4270  
 CCCCCTATTCCCTATTGTCTTACCTCGGTCTATTTTGCTTATCAGTGGACCTCACGAGGCACTCATAGGCATTT  
 4285 4295 4305 4315 4325 4335 4345  
 GAGTCTATGTGTCCTGTCCACATCCTCTGTAAAGGTGCAGAGAAAGTCCATGAGCAAGATGGAGCACTTCTAGTG  
 4360 4370 4380 4390 4400 4410 4420  
 GGTCCAAAGTCAGGGACACTATTTCAGCAATCTACAGTGCACAGGGCAGTTCCTCCCAACAGAGAATTACCTGTCTCTG  
 ApaI  
 4435 4445 4455 4465 4475 4485 4495  
 AATGTGGATCTGGCCCCCTTCCTTCCCCACTGTATAATGTGAAAACCTCTATGCTTTGTTCCCTTGTCTGCAAA  
 4510 4520 4530 4540 4550 4560 4570  
 ACAGGGATAATCCAGAACTGAGTTGTCCATGTAAAGTGCTTAGAACAGGGAGTGCTTGGCTTGGGGAGTGTCCAC  
 BS

APPROVED	O.G. FIG. none	
BY	CLARK	SUBCLASS
DRAFTSMAN	530	326

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 8861237

# FIG. 1A-6

4585 4595 4605 4615 4625 4635 4645  
 CTGCAGTCATTCATTATGCCCAGACAGGATGTTTCTTTATAGAAACGTGGAGGCCAGTTAGAACGACTCACCGCT  
 pMI+  
 PstI  
 4660 4670 4680 4690 4700 4710 4720  
 TCTCACCAC TGCCCATGTTTGTGTGTGTTCAGGTCCACTTCATCAACCCGGAAACGGTGCCCAAGCCCTGCT  
 PflMI  
 4735 4745 4755 4765 4775 4785 4795  
 GTGCGCCACGCAGCTCAATGCCATCTCCGTCTCTACTTCGATGACAGCTCCAACGTCATCCTGAAGAAATACA  
 4810 4820 4830 4840  
 GAAACATGGTGTCGGGCCTGTGGCTGCCACTAGCTCCTCCGA

APPROVED	O.G. FIG. none	
BY	CLASS	SUBCLASS
OR. FISHMAN	530	326

17621988  
 351

APPROVED BY O.G. FIG. none  
CLASS SUBCLASS  
DRAFTSMAN 530 326

FIG. 1B

CONSENSUS PROBE 20 30 40 50 60 70  
GATCCTAATGGGCTGTACGTGGACTTCCAGCGCGACGTGGGCTGGGACGACTGGATCATGCCCCCGTCG  
\*\*  
TGTAAGAAGCAGAGCTGTATGTCAGCTTCCGAGACCTGGGCTGGCAGGACTGGATCATGCGCCTGAAG  
OP4 28 38 48 58 68 78 88  
80 90 100 110 120 130 140  
ACTTCGACGCCCTACTACTGCTCCGGAGCCTGCCAGTTCCCTCTGCGGATCACTTCAACAGCACCAACCA  
\*\* \*\* \*\*\*\*\* \*\* \*\* \*\*\*\*\* \*\* \*\* \*\*\*\*\* \*\*  
GCTACGCGCGCTACTACTGTGAGGGGAGTGTGCCTTCCCTCTGAACTCCTACATGAACGCCCAACCA  
98 108 118 128 138 148 158  
150 160 170 180 190 200 210  
CGCCGTGGTGCAGACCCCTGGTGAACAACATGAACCCCGCAAGGTACCCAGCCCTGCTGCGTGCCACC  
\*\*\*\* \*\*\*\*\* \*\* \*\* \*\*\*\*\* \*\* \*\* \*\*\*\*\* \*\*  
CGCCATCGTGCAGACGCTGGTCCACTTTCATCAACCCGGAACGGTGCCCCAGCCCTGCTGTGCGCCCCACG  
168 178 188 198 208 218 228  
220 230 240 250 260 270 280  
GAGCTGTCCGCCATCAGCATGCTGTACCTGGACGAGAAATTCACCCGTGGTGTGAAGAACTACCAGGAGA  
\*\*\*\* \*\*\*\*\* \*\* \*\* \*\*\*\*\* \*\* \*\* \*\*\*\*\* \*\*  
CAGCTCAATGCCCATCTCCGTCCTCTACTTCGATGACAGCTCCCAACGTCATCCTGAAGAAATACAGAAACA  
238 248 258 268 278 288 298  
290 300 310  
TGACCGTGGTGGGCTGCGGCTGCCGCTAACTGCA  
\*\* \*\* \*\*\*\*\* \*\* \*\*  
TGGTGTCCGGGCGCTGTGGCTGCCACTAGCTCCT  
308 318 328

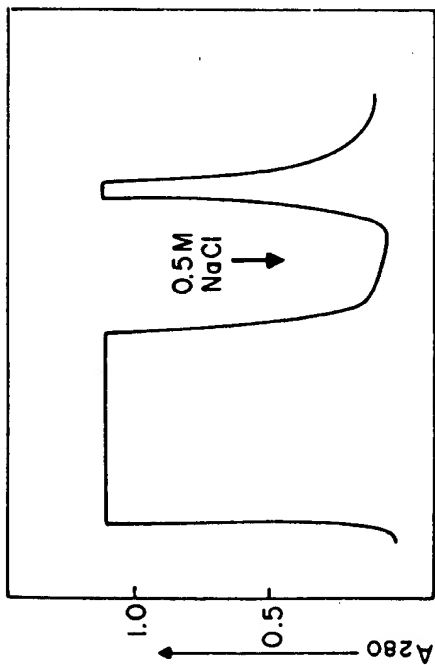


FIG. 2A

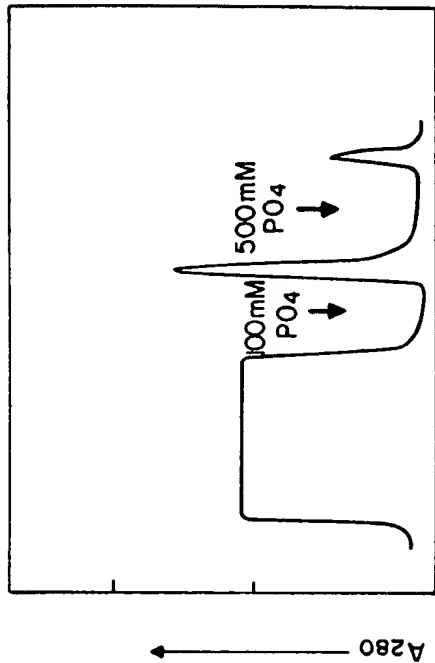


FIG. 2B

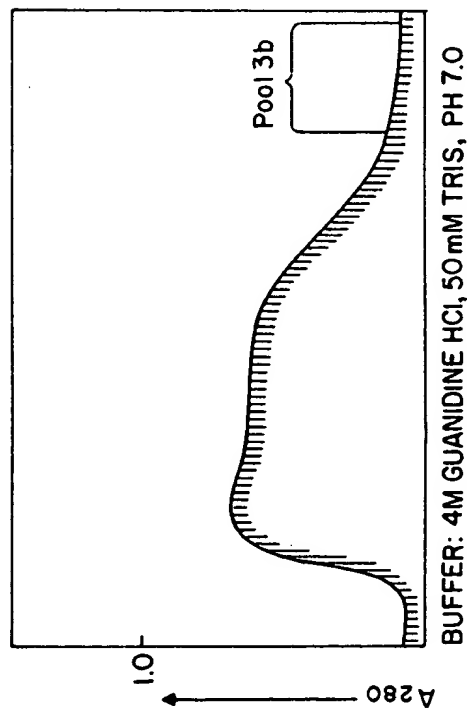
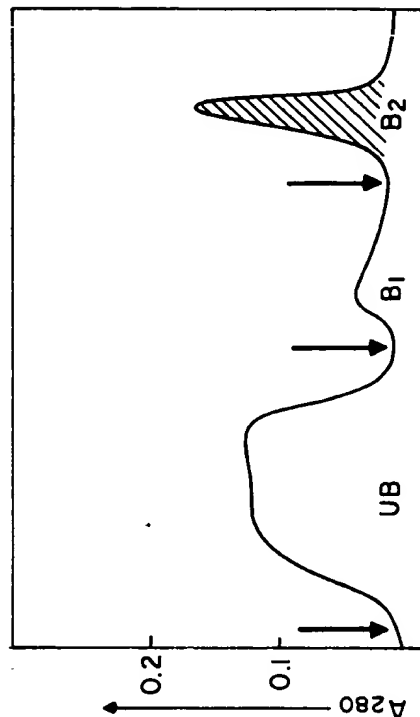


FIG. 2C



UB: 6M UREA 50mM TRIS 0.1M NaCl PH 7.0  
 B1: 6M UREA 50mM TRIS 0.5M NaCl PH 7.0  
 B2: 6M UREA 50mM TRIS 0.5M NaCl PH 7.0

FIG. 2D



FIG. 3A FIG. 3B



FIG. 4A FIG. 4B

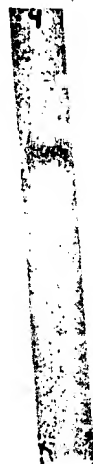
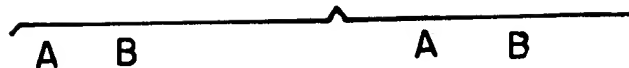


FIG. 5



1

2

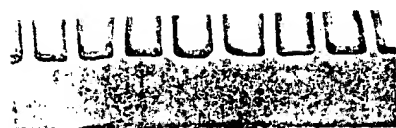
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FIG. 6 A FIG. 6 B FIG. 6 C FIG. 6 D FIG. 6 E



FIG. 15



— NON-REDUCIBLE 30K

— 18K SUBUNIT

— 16K SUBUNIT

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APPROVED	BY	CLASS	SUBCLASS
		570	326
O.G. FIG. 7A			

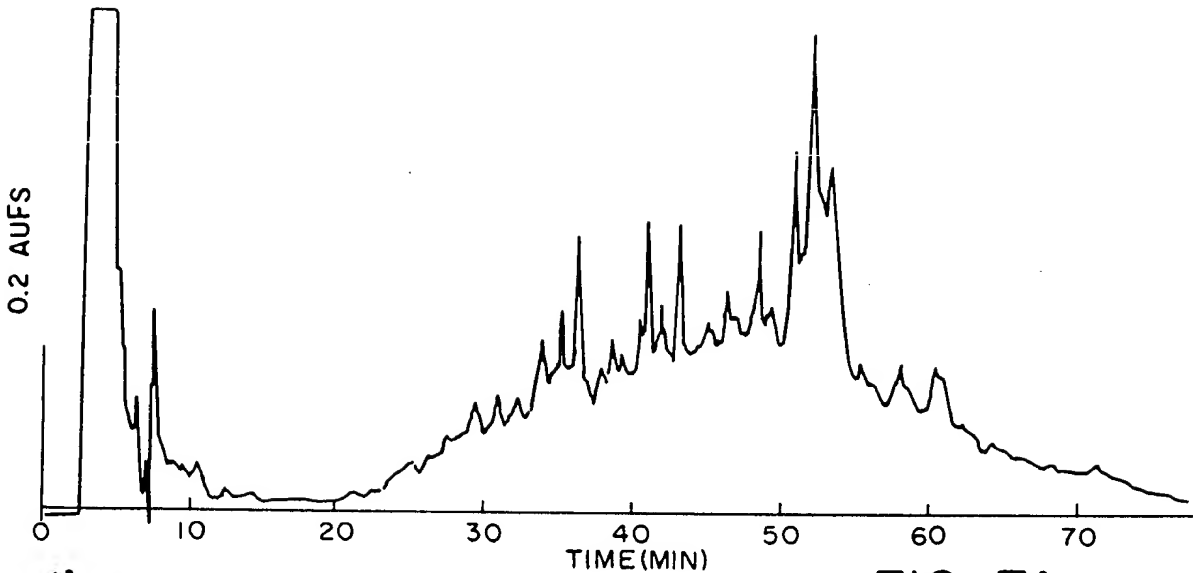


FIG. 7A

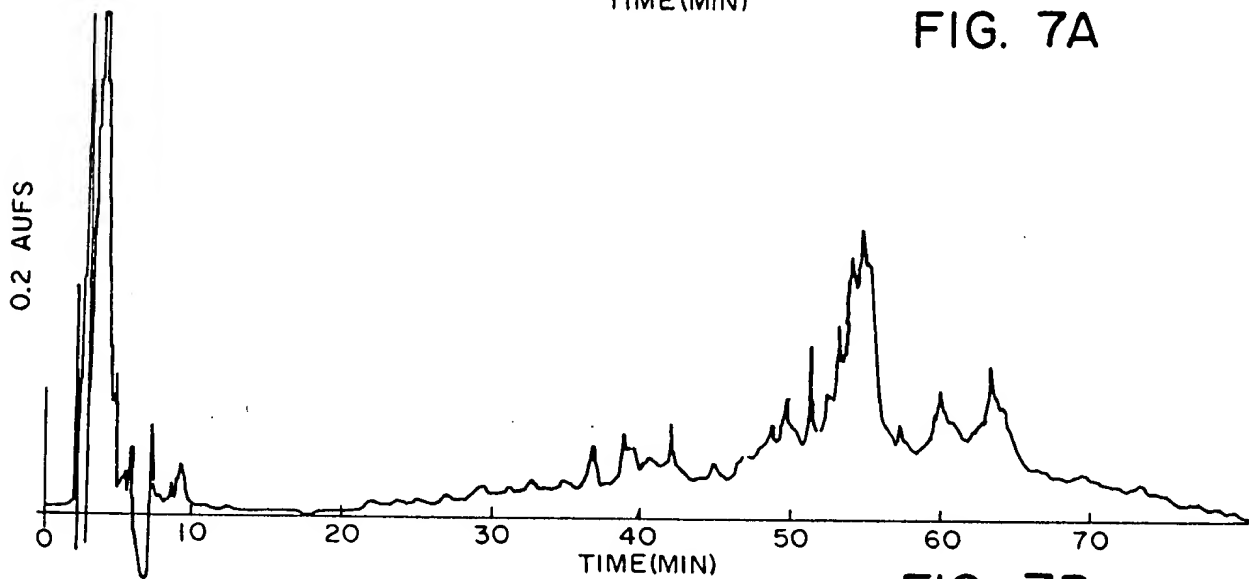


FIG. 7B

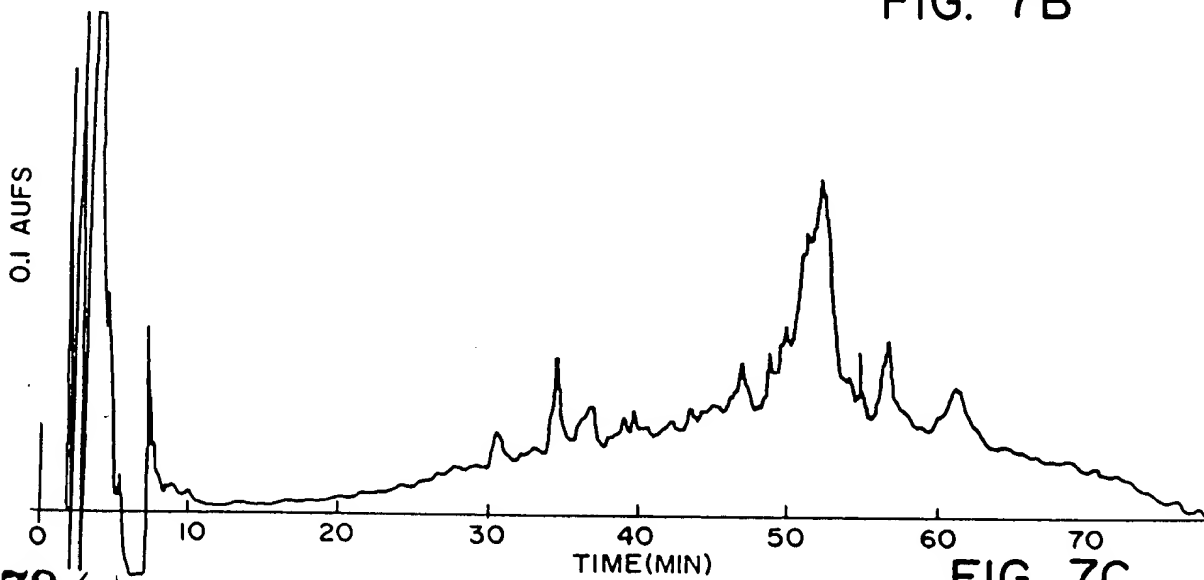


FIG. 7C

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APPROVED BY: O.G. FIG. 400  
 CLASS: 530  
 SUBCLASS: 326  
 DRAFTSMAN:

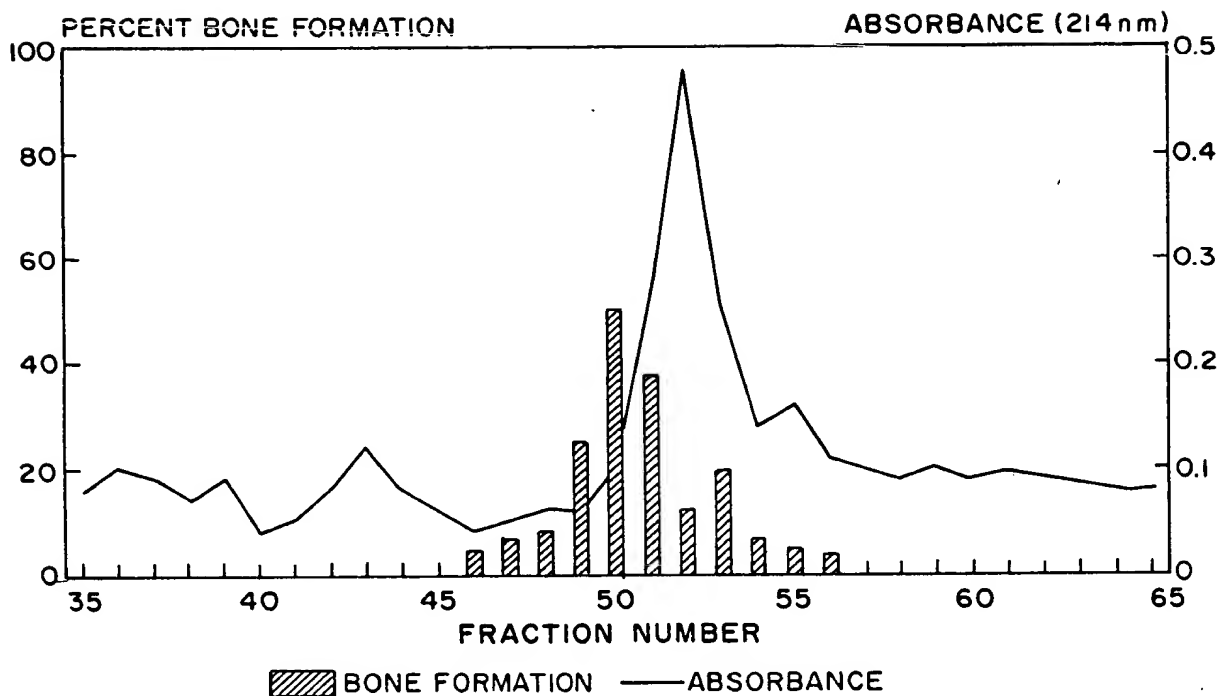


FIG. 8

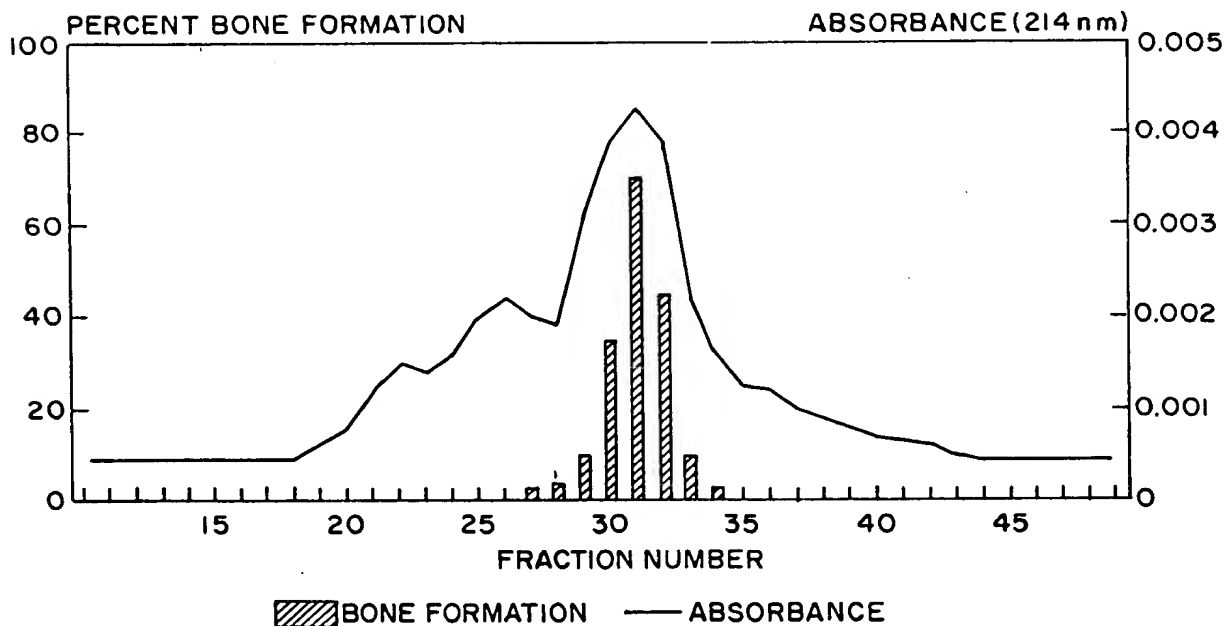


FIG. 9

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PURIFICATION SCHEME

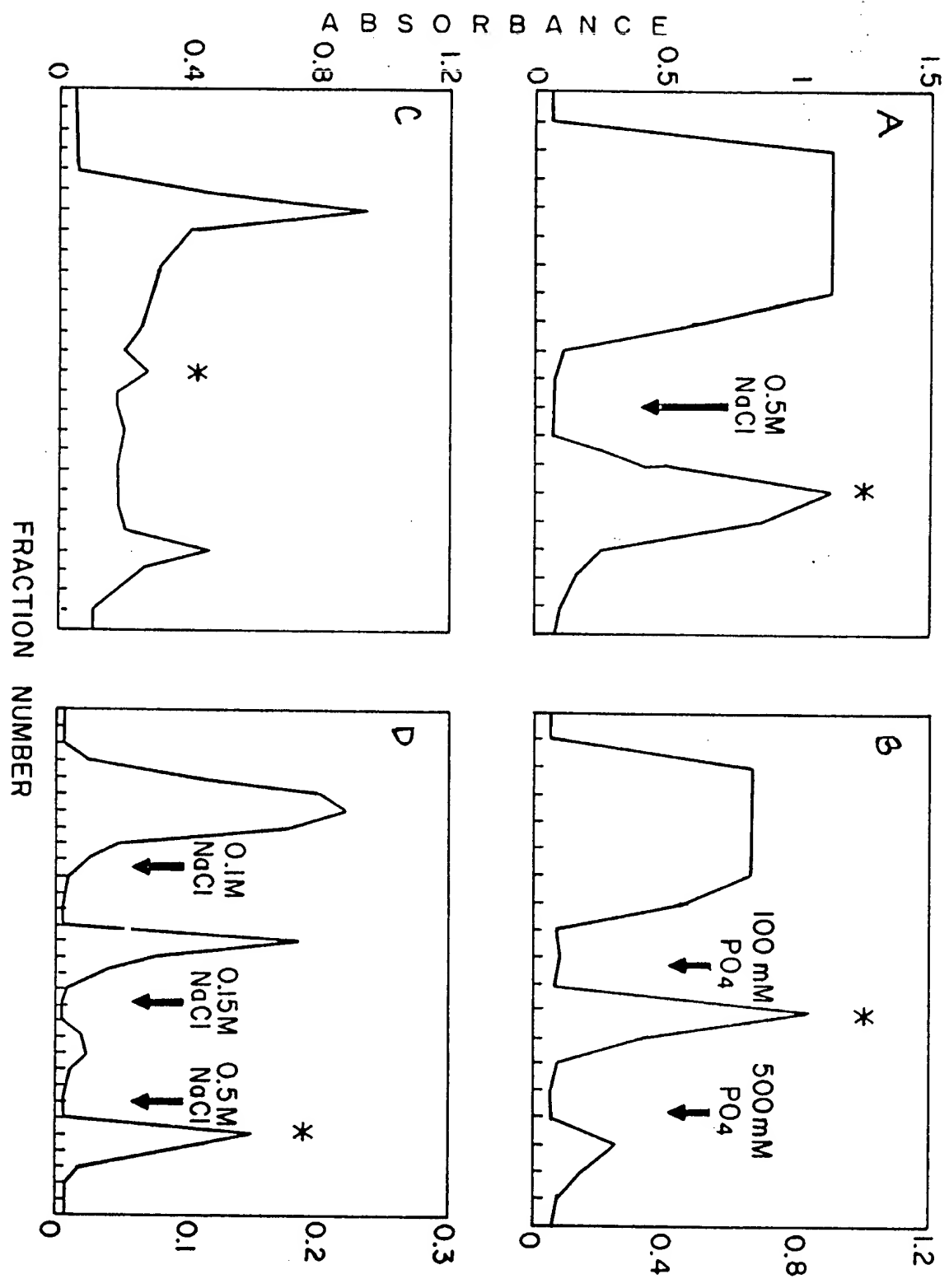


FIG. 10

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DOSE CURVES  
PERCENT BONE FORMATION

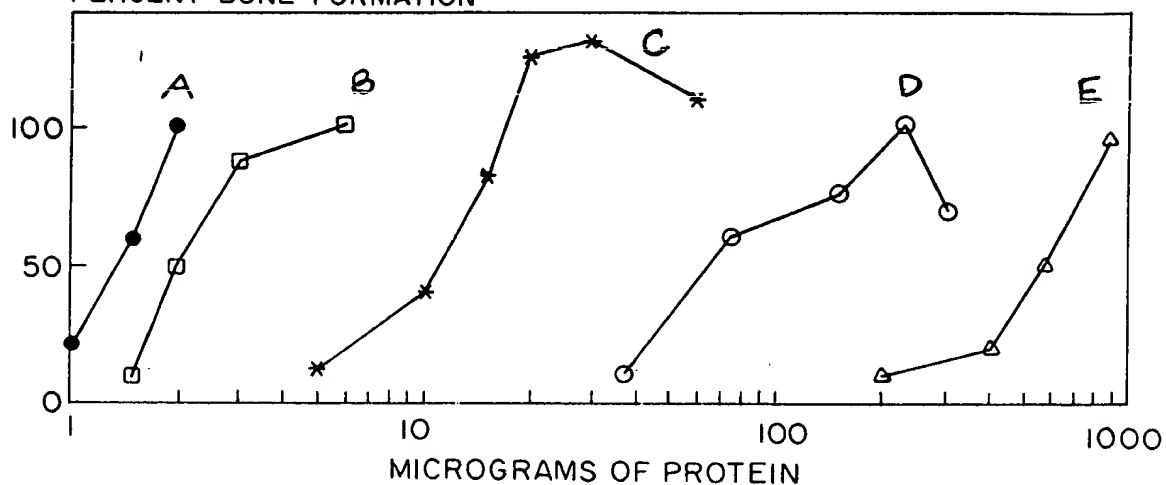


FIG. 11

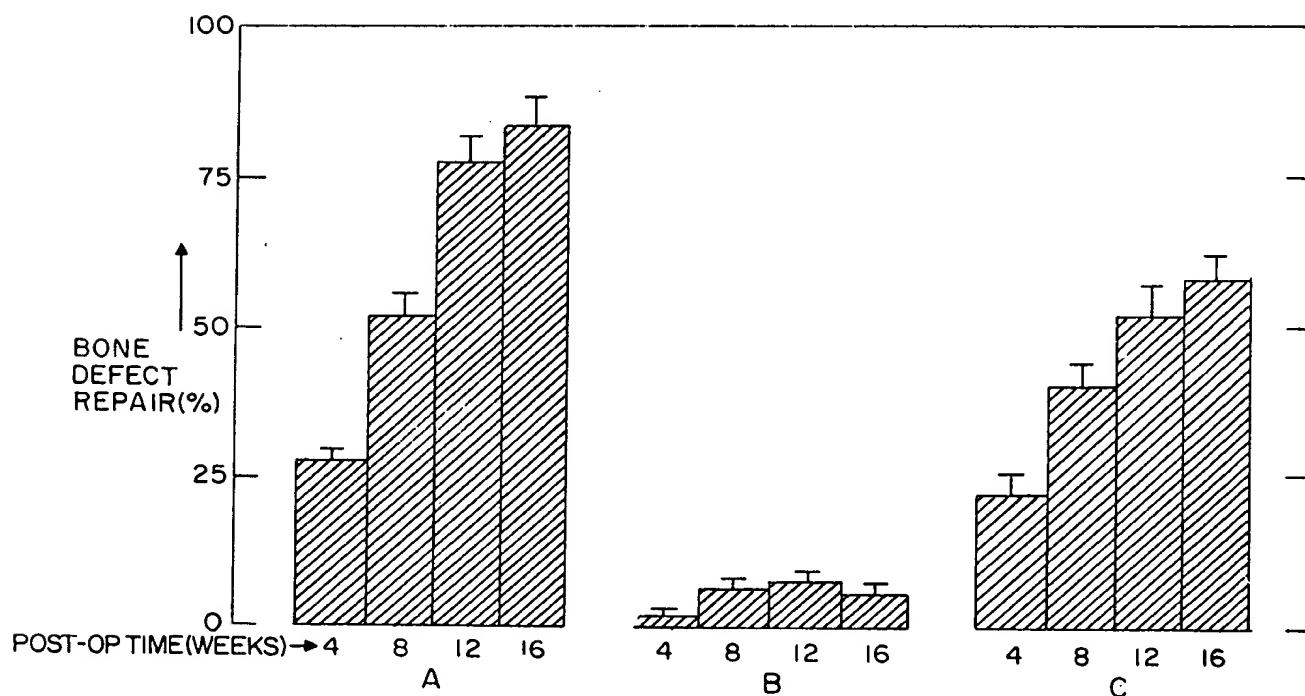


FIG. 12

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886123/11

APPROVED	O.G. FIGONE	CLASS	SUBCLASS
SY		030	326
DRAFTSMAN			

# FIG. 13

```

      10      20      30      40      50
GATCCTAATGGGCTGTACGTGGACTTCCAGCGCGACGTGGGCTGGGACGA
D P N G L Y V D F Q R D V G W D D

      60      70      80      90      100
CTGGATCATCGCCCCCGTCGACTTCGACGCCTACTACTGCTCCGGAGCCT
W I I A P V D F D A Y Y C S G A

      110     120     130     140     150
GCCAGTTCCCCTCTGCGGATCACTTCAACAGCACCAACCACGCCGTGGTG
C Q F P S A D H F N S T N H A V V

      160     170     180     190     200
CAGACCCTGGTGAACAACATGAACCCCGGCAAGGTACCCAAGCCCTGCTG
Q T L V N N M N P G K V P K P C C

      210     220     230     240     250
CGTGCCCAACCGAGCTGTCCGCCATCAGCATGCTGTACCTGGACGAGAATT
V P T E L S A I S M L Y L D E N

      260     270     280     290     300
CCACCGTGCTGAAGAACTACCAGGAGATGACCGTGGTGGGCTGCGGC
S T V V L K N Y Q E M T V V G C G

      310
TGCCGCTAACTGCAG
C R *

```

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SDS GEL ELUTION OF OSTEOGENIC ACTIVITY  
CALCIUM CONTENT (ug/mg tissue)

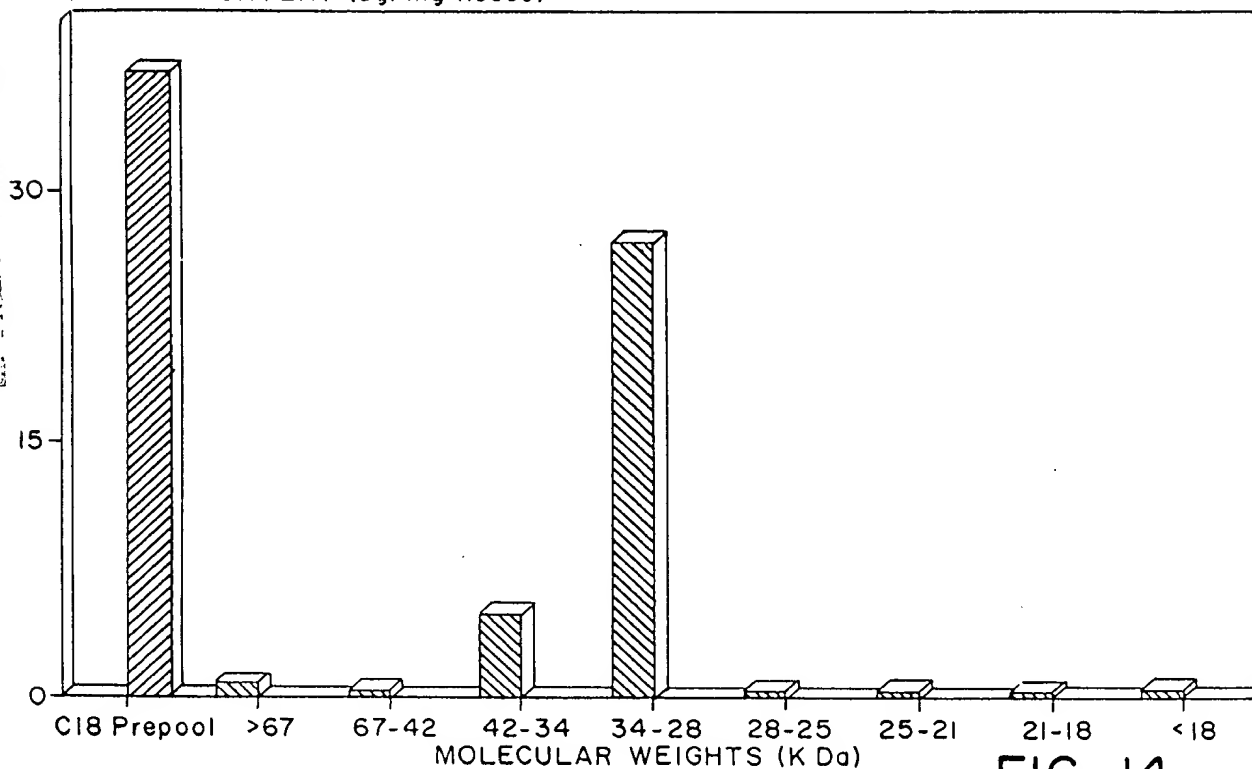


FIG. 14

ALKALINE PHOSPHATASE (U/mg protein)

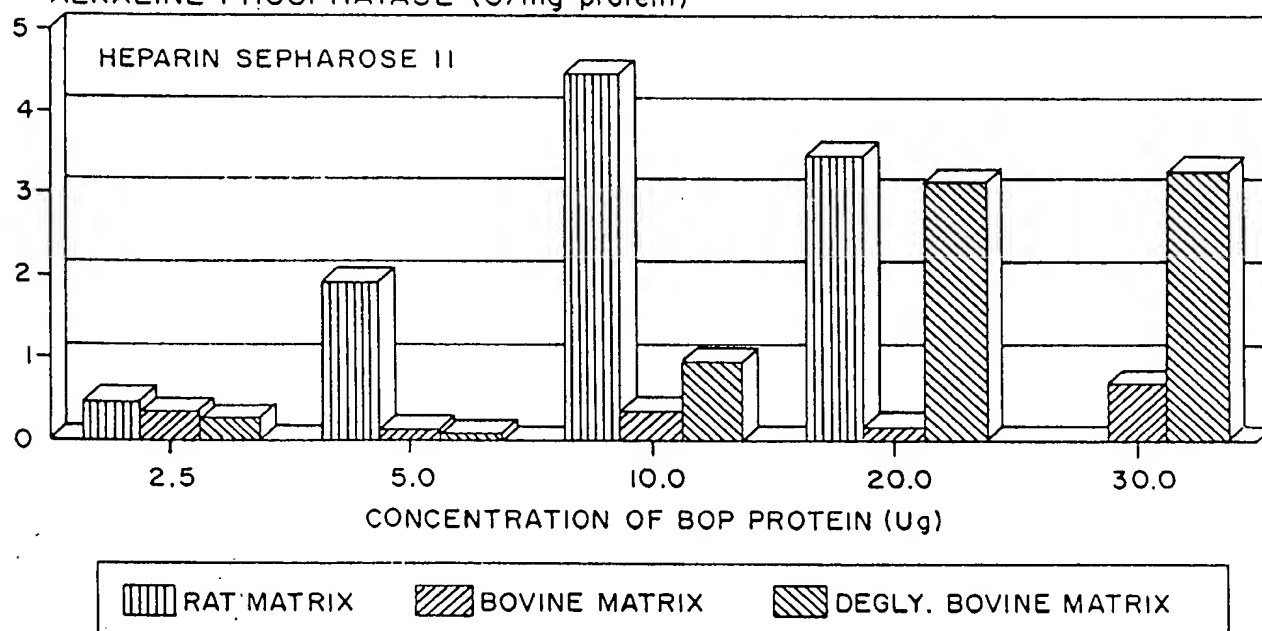


FIG. 19

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APPROVED BY: *O.G. Figueira*  
CLASS: 530  
SUBCLASS: 326  
DRAWN BY: [blank]

HPLC PROFILE  
ENDO ASP-N DIGEST - PREPOOL 16K OP SUBUNIT

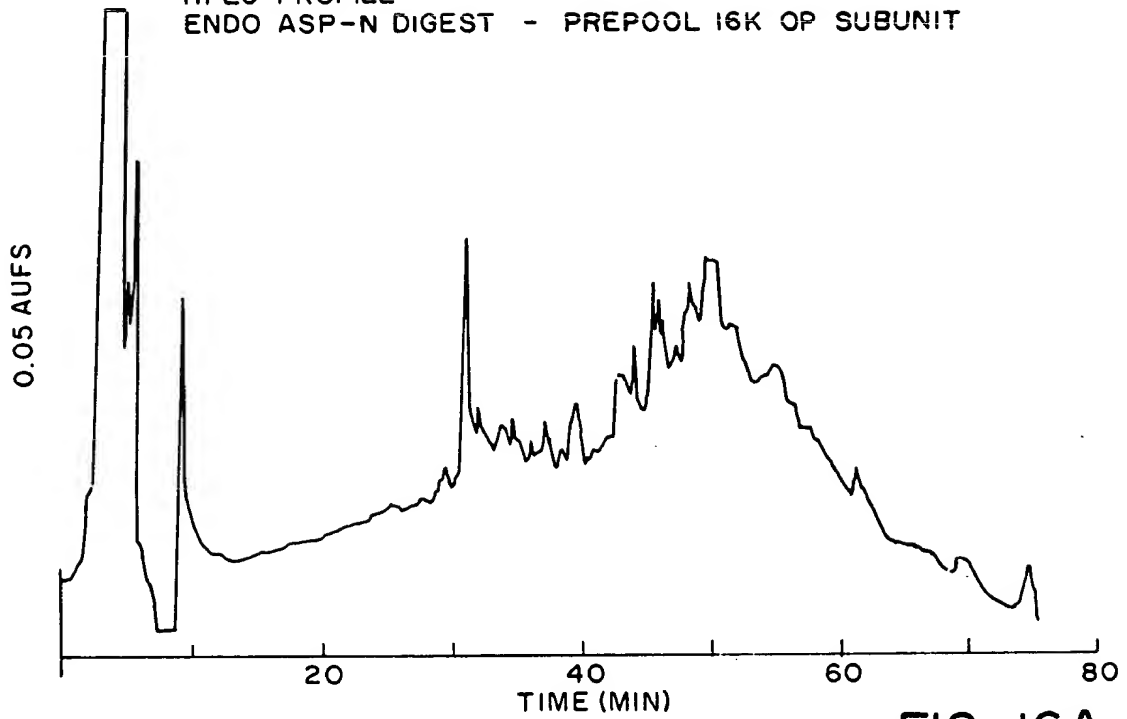


FIG. 16A

HPLC PROFILE  
ENDO ASP-N DIGEST - PREPOOL 18K OP SUBUNIT

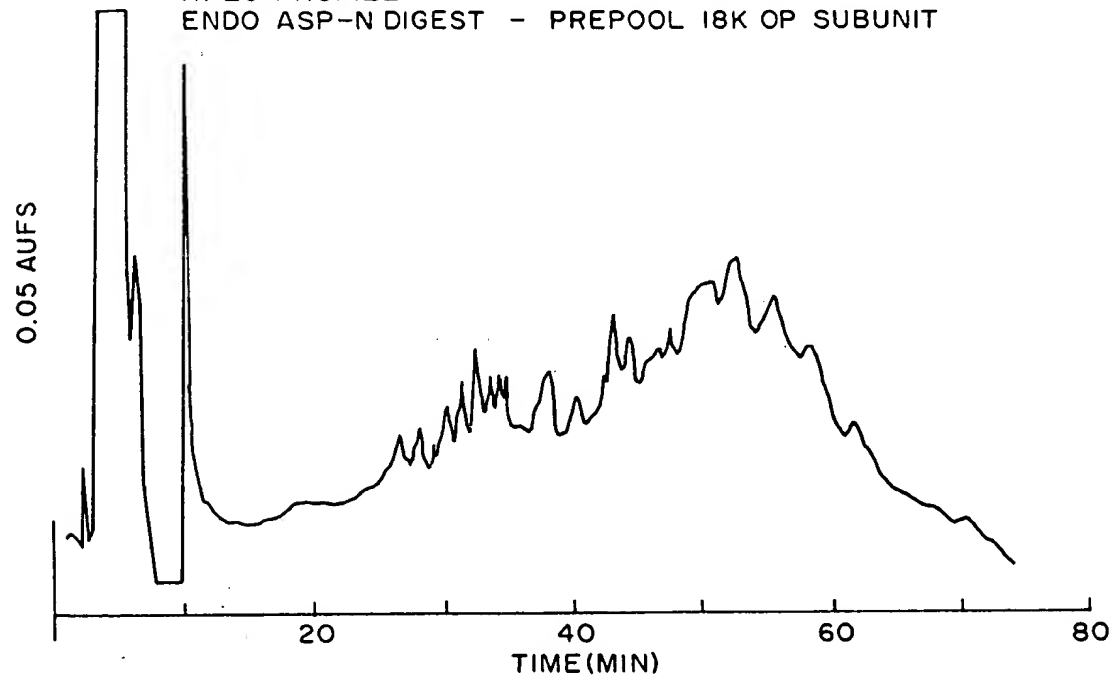


FIG. 16B

886129/10  
*347*

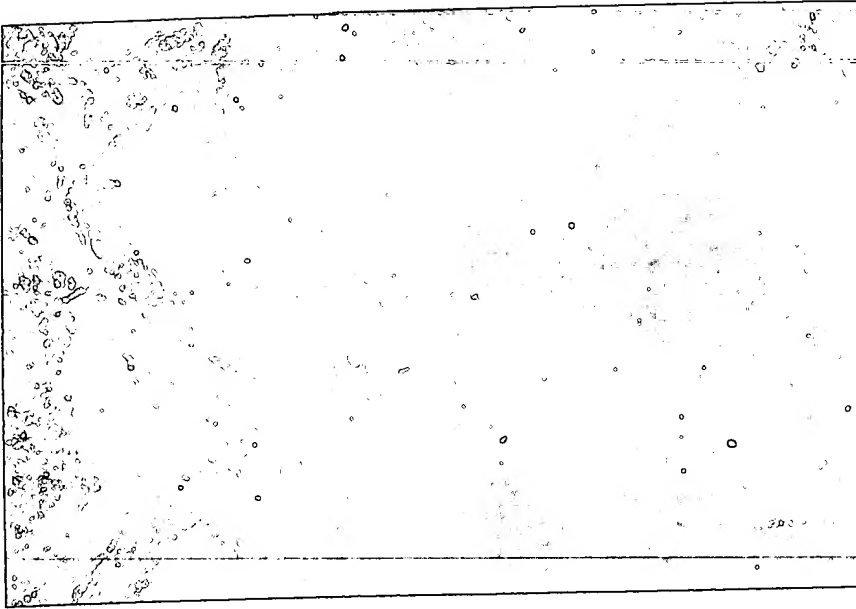


FIG. 17A

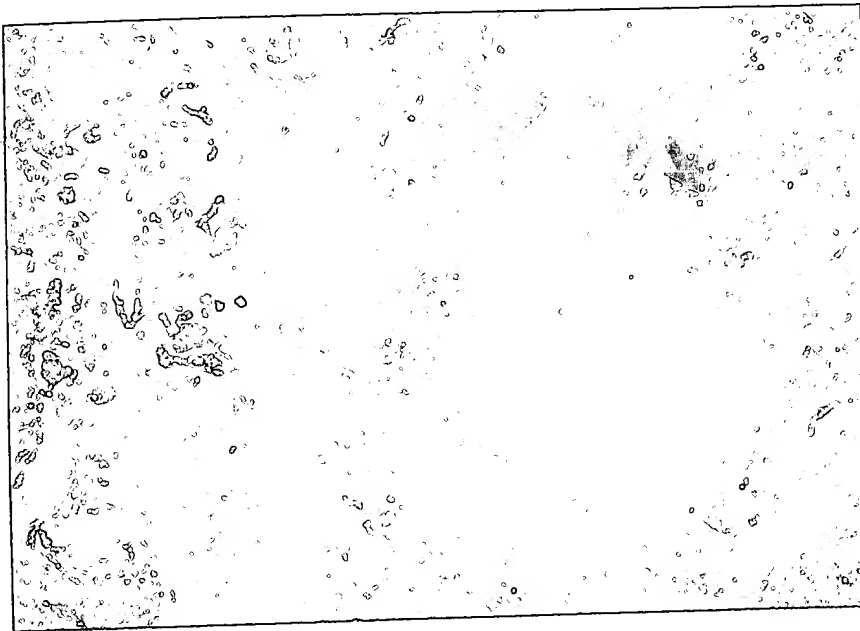


FIG. 17 B

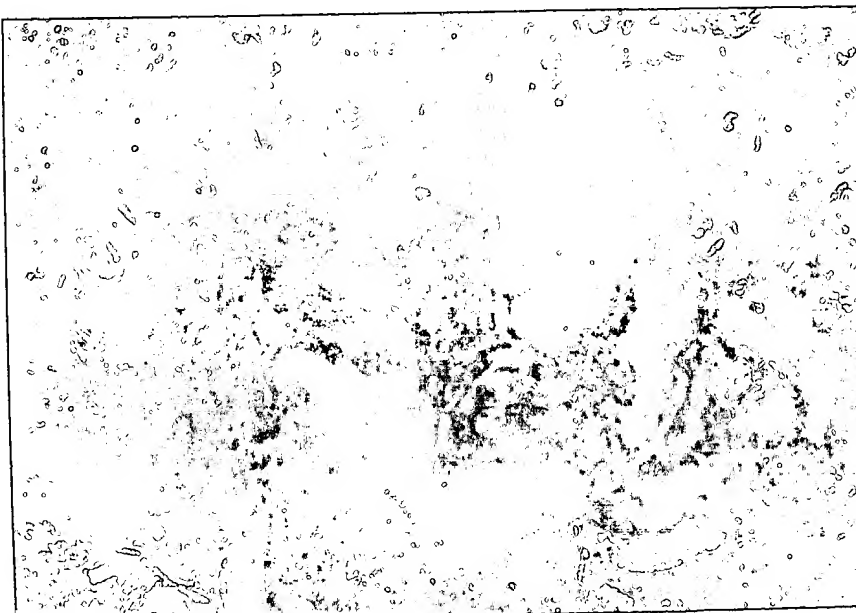


FIG. 17C

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11/521988





APPROVED	O.G. FIG. 204
BY	CLASS
ORATSIAN	SUBCLASS
530	326

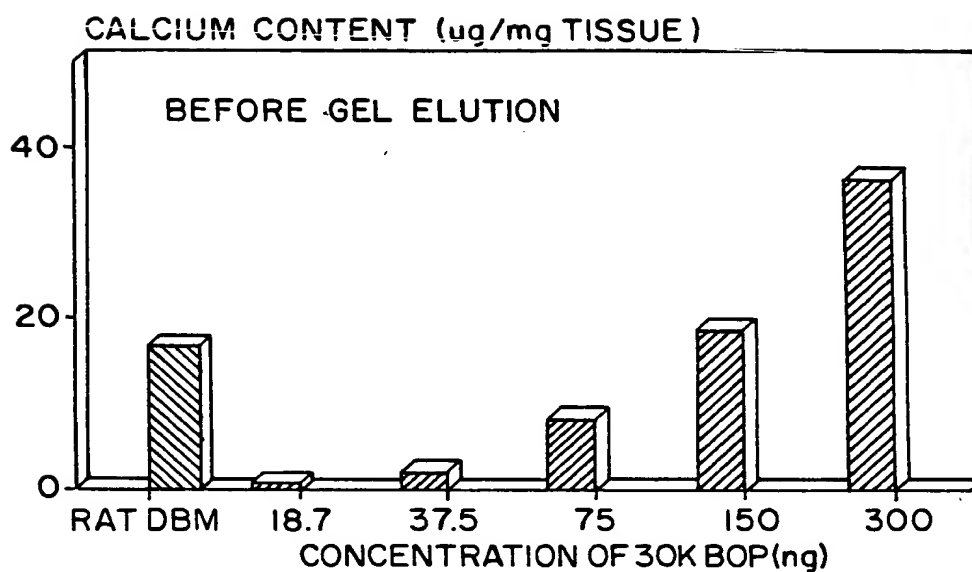
S	L	L	P	G	A	Q	P	C	C	A	L	P	G	T	M	R	P	L	H	V	R	T	T	S	D				
Q	A	R	G	A	L	A	R	P	P	C	C	V	P	T	A	Y	A	G	K	L	L	I	S	L	S	E	E	R	
T	T	L	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:		
Q	T	I	L	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:		
H	N	P	G	E	A	S	A	S	A	S	P	Q	A	D	L	E	P	L	P	T	I	V	Y	Y	V	G	R	K	
R	G	L	N	P	G	T	K	V	N	S	C	C	I	P	T	K	L	S	T	M	S	M	L	Y	F	D	D	E	Y
R	G	H	S	P	F	A	N	L	K	S	C	C	V	P	T	K	L	R	P	M	S	M	L	Y	Y	D	D	G	Q
A	V	G	V	V	P	G	I	P	E	P	C	C	V	P	E	K	M	S	S	L	S	I	L	F	F	D	E	N	K
:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	
S	V	N	S	K	I	P	K	A	C	C	V	P	T	E	L	S	A	I	S	M	L	Y	L	D	E	N	E	E	
F	I	N	P	E	T	V	P	K	A	C	C	A	P	T	Q	L	N	A	I	S	V	L	Y	F	D	D	S	S	
N	N	N	P	G	K	V	P	K	A	C	C	V	P	T	Q	L	D	S	V	A	M	L	Y	L	N	D	Q	S	
S	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	
S	S	V	N	S	K	I	P	K	A	C	C	V	P	T	E	L	S	A	I	S	M	L	Y	L	D	E	N	E	
S	S	V	N	S	K	I	P	K	A	C	C	V	P	T	E	L	S	A	I	S	M	L	Y	L	D	E	N	E	
N	M	N	P	G	K	V	P	K	P	C	C	V	P	T	E	L	S	A	I	S	M	L	Y	L	D	E	N	E	
N	M	N	P	G	K	V	P	K	P	C	C	V	P	T	E	L	S	A	I	S	M	L	Y	L	D	E	N	E	
N	M	N	P	G	K	V	P	K	P	C	C	V	P	T	E	L	S	A	I	S	M	L	Y	L	D	E	N	E	

FIG. 18-3

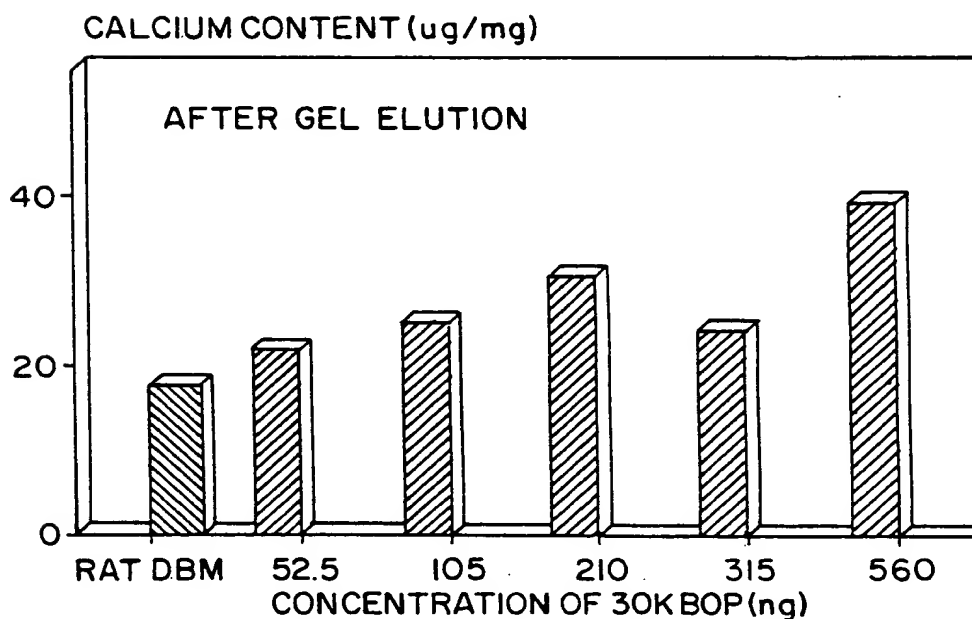
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APPROVED	BY	DRAFTSMAN
O.G. FIG. 20A	CLASS	SUBCLASS
	530	326



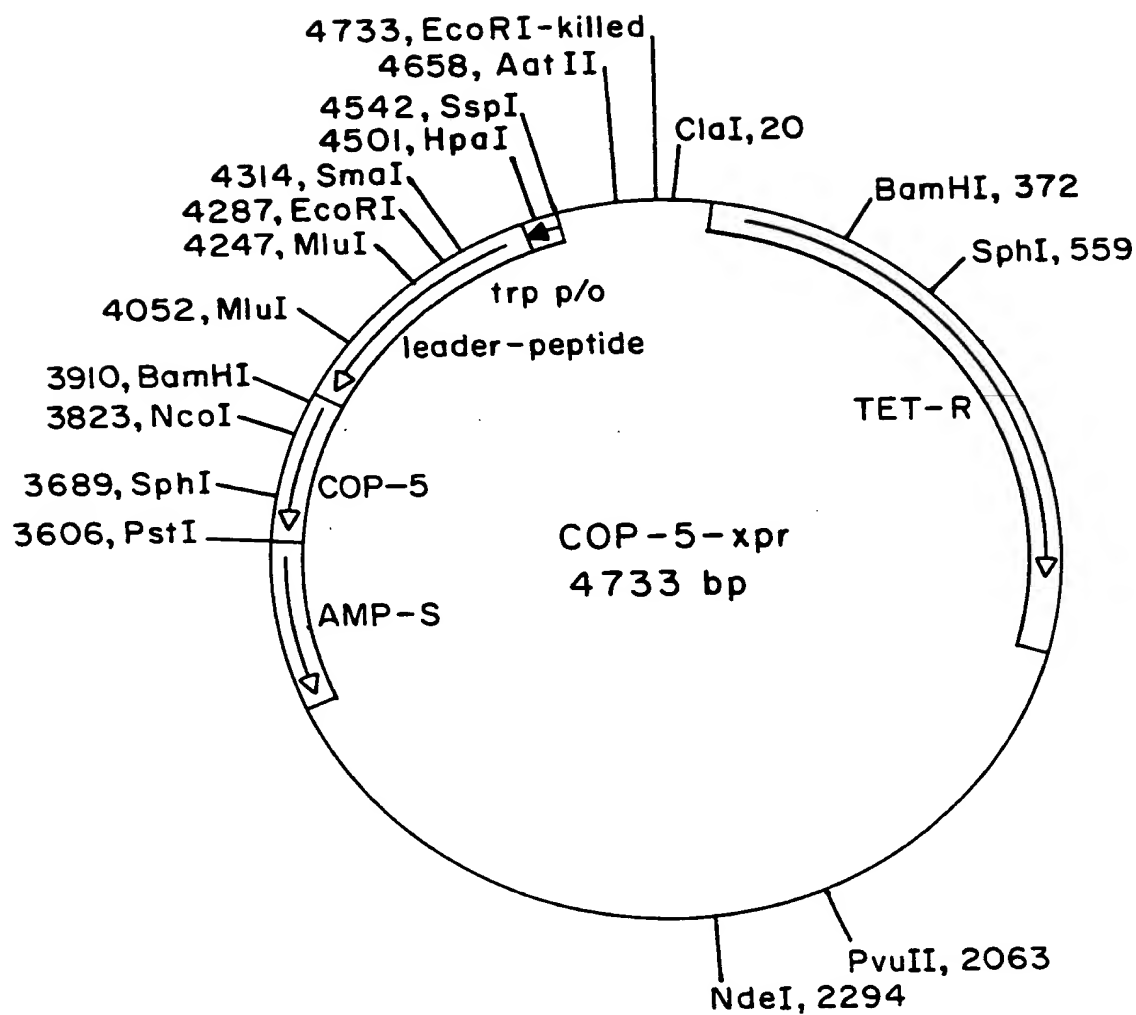
**FIG. 20A**



**FIG. 20B**

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APPROVED	BY	CLASS	SUBCLASS
0.6. FIG. 204	530	326	
DRAFTSMAN			



**FIG. 21A**

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# COP-5 fusion protein

10 20 30 40 50  
 ATGAAAGCAATTTTCGTACTGAAAGGTTCACTGGACAGAGATCTGGACTC  
 M K A I F V L K G S L D R D L D S  
 BglII

60 70 80 90 100  
 TCGTCTGGATCTGGACGTTTCGTACCGACCACAAAGACCTGTCTGATCACC  
 R L D L D V R T D H K D L S D H

110 120 130 140 150  
 TGGTTCTGGTCGACCTGGCTCGTAACGACCTGGCTCGTATCGTTACTCCC  
 L V L V D L A R N D L A R I V T P  
 SalI Sma

160 170 180 190 200  
 GGGTCTCGTTACGTTGCGGATCTGGAATTCATGGCTGACAACAAATTCAA  
 G S R Y V A D L E F M A D N K F N  
 I EcoRI

210 220 230 240 250  
 CAAGGAACAGCAGAACGCGTTCTACGAGATCTTGACCTGCCGAACCTGA  
 K E Q Q N A F Y E I L H L P N L  
 MluI BglII BspMI+

260 270 280 290 300  
 ACGAAGAGCAGCGTAACGGCTTCATCCAAAGCTTGAAGGATGAGCCCTCT  
 N E E Q R N G F I Q S L K D E P S  
 HindIII

310 320 330 340 350  
 CAGTCTGCCAATCTGCTAGCGGATGCCAAGAACTGAACGATGCGCAGGC  
 Q S A N L L A D A K K L N D A Q A  
 NheI FspI

360 370 380 390 400  
 ACCGAAATCGGATCAGGGGCAATTCATGGCTGACAACAAATTCAACAAGG  
 P K S D Q G Q F M A D N K F N K

410 420 430 440 450  
 AACAGCAGAACGCGTTCTACGAGATCTTGACCTGCCGAACCTGAACGAA  
 E Q Q N A F Y E I L H L P N L N E  
 MluI BglII BspMI+

460 470 480 490 500  
 GAGCAGCGTAACGGCTTCATCCAAAGCTTGAAGGATGAGCCCTCTCAGTC  
 E Q R N G F I Q S L K D E P S Q S  
 HindIII

FIG. 21B-1

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O.G. FIG. 204	
CLASS	SUBCLASS
530	326
APPROVED	BY
	DRAFTSMAN

510 520 530 540 550  
TGC GAATCTGCTAGCGGATGCCAAGAACTGAACGATGCGCAGGCACCGA  
A N L L A D A K K L N D A Q A P  
NheI FspI

560 570 580 590 600  
AGGATCCTAATGGGCTGTACGTCGACTTCAGCGACGTGGGCTGGGACGAC  
K D P N G L Y V D F S D V G W D D  
BamHI SalI

610 620 630 640 650  
TGGATTGTGGCCCCACCAGGCTACCAGGCCTTCTACTGCCATGGCGAATG  
W I V A P P G Y Q A F Y C H G E C  
StuI NcoI BsmI+

660 670 680 690 700  
CCCTTTCCCGCTAGCGGATCACTTCAACAGCACCAACCACGCCGTGGTGC  
P F P L A D H F N S T N H A V V  
NheI DraIII  
PflMI

710 720 730 740 750  
AGACCCTGGTGAAGTCTGTCAACTCCAAGATCCCTAAGGCTTGCTGCGTG  
Q T L V N S V N S K I P K A C C V  
MstII

760 770 780 790 800  
CCCACCGAGCTGTCCGCCATCAGCATGCTGTACCTGGACGAGAATGAGAA  
P T E L S A I S M L Y L D E N E K  
SphI

810 820 830 840 850  
GGTGGTGCTGAAGAACTACCAGGAGATGGTAGTAGAGGGCTGCGGCTGCC  
V V L K N Y Q E M V V E G C G C  
PflMI

860  
GCTAACTGCAG  
R \*  
PstI

*FIG. 21B-2*

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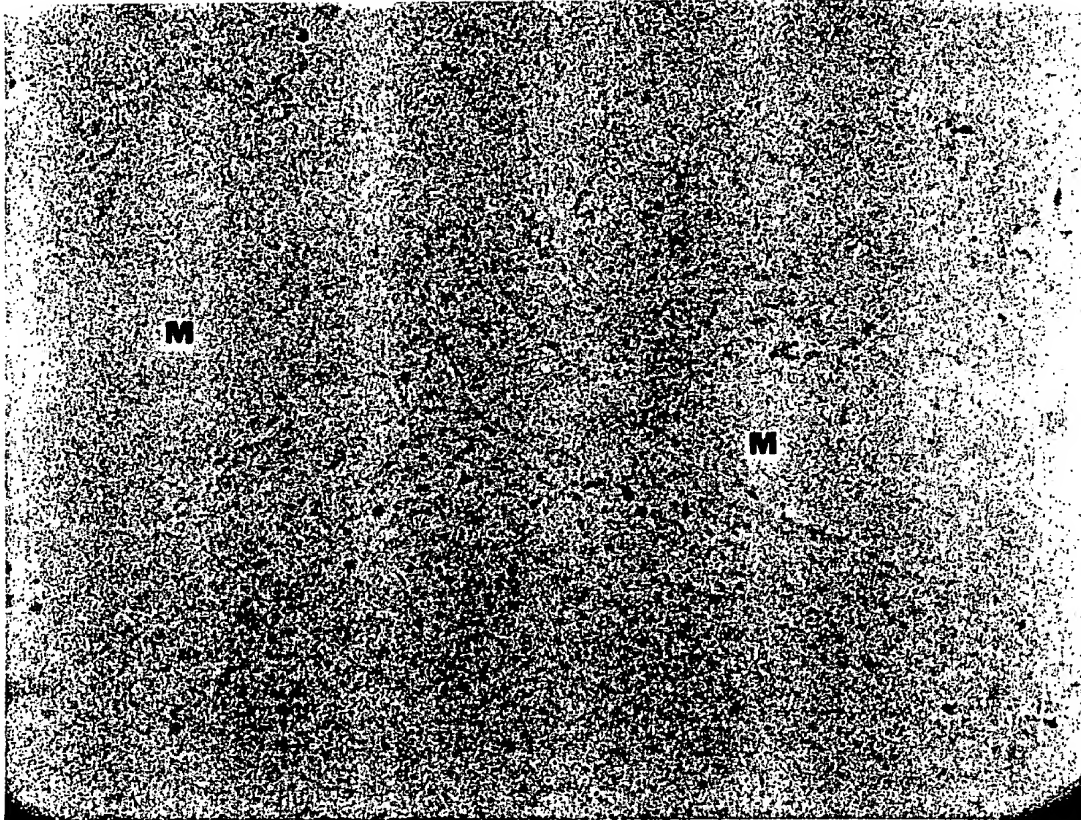


FIG. 22A

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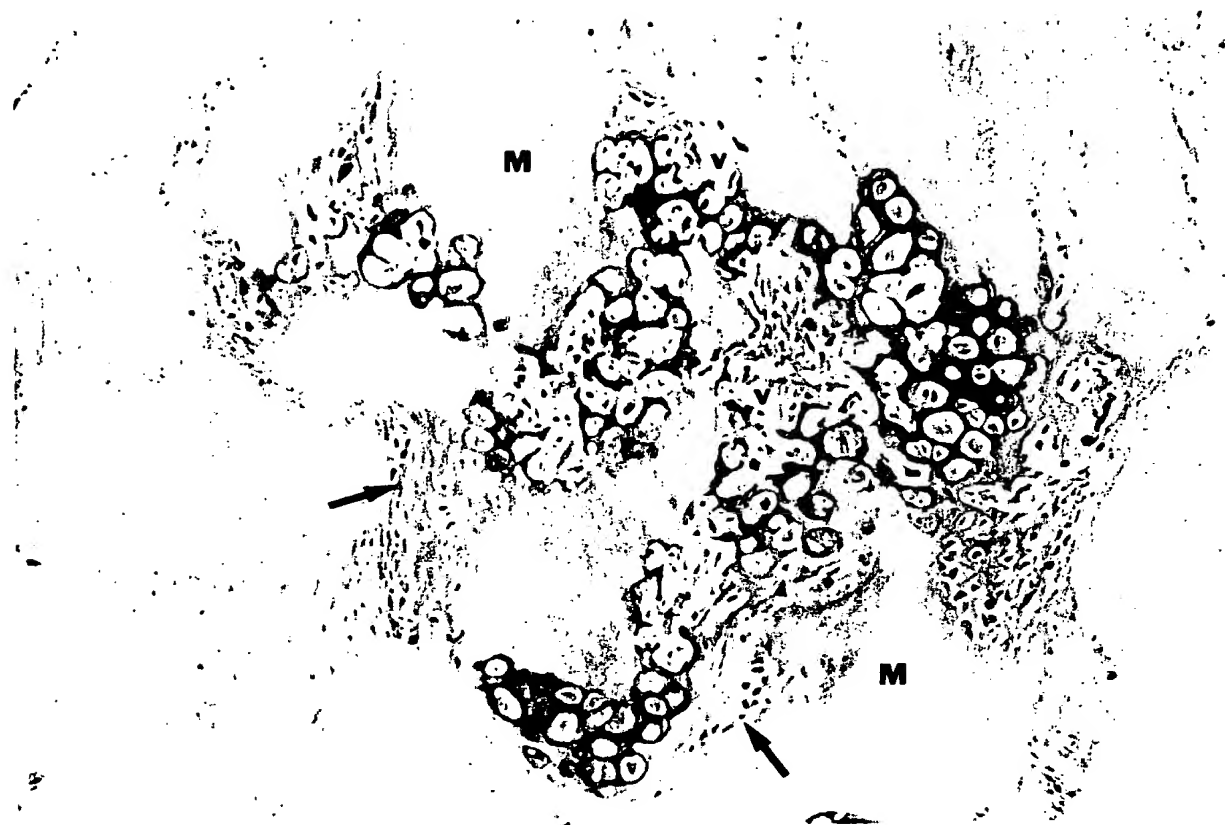


FIG. 22B

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